

# HMP2 IBD metagenome and metatranscriptome profiling

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Updated date: Sep 8, 2021

 An abbreviated version of this protocol was published in eLIFE in May 2021

Integrating taxonomic, functional, and strain-level profiling of diverse microbial communities with bioBakery 3

DOI: 10.7554/eLife.65088

## Detailed protocol

Please check <https://forum.biobakery.org/> for answers to questions about bioBakery tools. If you don't already see your question there, please feel free to start a new thread. Thanks!

**How to cite:** (Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Huttenhower, C. , Franzosa, E. and Segata, N. (2021). HMP2 IBD metagenome and metatranscriptome profiling. Bio-protocol Preprint. [bio-protocol.org/prep1365](https://bio-protocol.org/prep1365).
2. Beghini, F., McIver, L. J., Blanco-Míguez, A., Dubois, L., Asnicar, F., Maharjan, S., Mailyan, A., Manghi, P., Scholz, M., Thomas, A. M., Valles-Colomer, M., Weingart, G., Zhang, Y., Zolfo, M., Huttenhower, C., Franzosa, E. A. and Segata, N.(2021). Integrating taxonomic, functional, and strain-level profiling of diverse microbial communities with bioBakery 3. eLIFE. DOI: [10.7554/eLife.65088](https://doi.org/10.7554/eLife.65088)

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